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TECH CENTER 1600/2900



1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/782,130

DATE: 11/04/2002

TIME: 13:28:08

Input Set : A:\seq1st.txt

Output Set: N:\CRF4\11042002\I782130.raw

3 <110> APPLICANT: KNAUF, VIC C.
4 KRIDL, JEAN C.
6 <120> TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR REGULATED TRANSCRIPTION
7 AND EXPRESSION OF HETEROLOGOUS GENES
9 <130> FILE REFERENCE: 16518.052
11 <140> CURRENT APPLICATION NUMBER: US 09/782,130
12 <141> CURRENT FILING DATE: 2001-02-12
14 <150> PRIOR APPLICATION NUMBER: US 09/232,861
15 <151> PRIOR FILING DATE: 1999-01-15
17 <150> PRIOR APPLICATION NUMBER: US 08/812,665
18 <151> PRIOR FILING DATE: 1997-03-07
20 <150> PRIOR APPLICATION NUMBER: US 08/484,941
21 <151> PRIOR FILING DATE: 1995-06-07
23 <150> PRIOR APPLICATION NUMBER: US 08/105,852
24 <151> PRIOR FILING DATE: 1993-08-10
26 <150> PRIOR APPLICATION NUMBER: US 07/526,123
27 <151> PRIOR FILING DATE: 1990-05-21
29 <150> PRIOR APPLICATION NUMBER: US 07/267,685
30 <151> PRIOR FILING DATE: 1988-11-02
32 <150> PRIOR APPLICATION NUMBER: US 06/692,605
33 <151> PRIOR FILING DATE: 1985-01-17
35 <150> PRIOR APPLICATION NUMBER: US 07/582,241
36 <151> PRIOR FILING DATE: 1990-09-14
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39 <151> PRIOR FILING DATE: 1988-04-29
41 <150> PRIOR APPLICATION NUMBER: US 07/168,190
42 <151> PRIOR FILING DATE: 1988-03-15
44 <150> PRIOR APPLICATION NUMBER: US 07/054,369
45 <151> PRIOR FILING DATE: 1987-05-26
47 <150> PRIOR APPLICATION NUMBER: US 07/742,834
48 <151> PRIOR FILING DATE: 1991-08-08
50 <150> PRIOR APPLICATION NUMBER: US 07/550,804
51 <151> PRIOR FILING DATE: 1990-07-09
53 <150> PRIOR APPLICATION NUMBER: US 07/147,781
54 <151> PRIOR FILING DATE: 1988-01-25
56 <150> PRIOR APPLICATION NUMBER: US 07/078,538
57 <151> PRIOR FILING DATE: 1987-07-28
59 <150> PRIOR APPLICATION NUMBER: US 06/891,529
60 <151> PRIOR FILING DATE: 1986-07-31
62 <160> NUMBER OF SEQ ID NOS: 35
64 <170> SOFTWARE: PatentIn Ver. 2.1
66 <210> SEQ ID NO: 1
67 <211> LENGTH: 2200

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Input Set : A:\seq1st.txt

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68 <212> TYPE: DNA

69 <213> ORGANISM: Brassica napus

71 <400> SEQUENCE: 1

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74 ctaaagggtt ctggctgttt tcttggtcat atgattaact tctaaacttg tgtataaata 180
75 ttctctgaaa gtgcttcttt tggcatatgt aggttgggca aaaacgagga agattgcttc 240
76 tcaatttgga agatgatgaa cagccgaaga agaaaataag aataggcagt cctgctactc 300
77 aatggatctc agtctataac ggtcgtcgtc ccatgaaaca gaggtaacac attttttgca 360
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80 ctgtgtagca tcagcagcta atctctgggc tctcatcatg gatgctggaa ctggattcac 540
81 ttctcaagtt tatgagttgt caccggtctt cctacacaag gtaataatca gttgaagcaa 600
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83 gactggatta tggacaatg ggaaaagaac tactatataa gctccatagc gggttcagat 720
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85 atcaaaaagt cttcattgac ttatttatat acttggttg aattggtagg aactacttat 840
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87 aaagaagatt ttcattgtaac ctccatgaca actgctggta atcgttgggg tgtggtaatg 960
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89 attttccctt gatagtctaa tatgataaac tctgcgttgt gaaagggtgt ggagcttgac 1080
90 tttttgtacc caagcgatgg gatacatagg aggtgggaga atgggtatag aataacatca 1140
91 atggcagcaa ctgcgatca agcagcttct atattaagca taccaaagcg taagatgggtg 1200
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105 cacatcacat attagcatgc agatgcggac gatttgtcac tcacttcaaa cacctaaaag 2040
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112 <211> LENGTH: 4324

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118 <222> LOCATION: (2152)..(2703)

120 <220> FEATURE:

121 <221> NAME/KEY: modified_base

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Input Set : A:\seq1st.txt

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147 ttctctgaaa gtgcttcttt tggcatatgt aggttgggca aaaacgagga agattgcttc 240
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167 ccacaaagtc ttcattgact tttttatata cttgtttgta attgctagga actacttatt 840
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173 cgaggaactc tggcttctct gatcaggtag gtttttgtct cttattgtct ggtgttttta 1020
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212                                     Met Ala
213                                     1
215 aac aag ctc ttc ctc gtc tcg gca act ctc gcc ttg ttc ttc ctt ctc 2205
216 Asn Lys Leu Phe Leu Val Ser Ala Thr Leu Ala Leu Phe Phe Leu Leu
217           5           10           15
219 acc aat gcc tcc gtc tac agg acg gtt gtg gaa gtc gac gaa gat gat 2253
220 Thr Asn Ala Ser Val Tyr Arg Thr Val Val Glu Val Asp Glu Asp Asp
221       20           25           30
223 gcc aca aat cca gcc ggc cca ttt agg att cca aaa tgt agg aag gag 2301
224 Ala Thr Asn Pro Ala Gly Pro Phe Arg Ile Pro Lys Cys Arg Lys Glu
225   35           40           45           50
227 ttt cag caa gca caa cac ctg aaa gct tgc caa caa tgg ctc cac aag 2349
228 Phe Gln Gln Ala Gln His Leu Lys Ala Cys Gln Gln Trp Leu His Lys
229           55           60           65
231 cag gca atg cag tcc ggt agt ggt cca agc tgg acc ctc gat ggt gag 2397
232 Gln Ala Met Gln Ser Gly Ser Gly Pro Ser Trp Thr Leu Asp Gly Glu
233           70           75           80
235 ttt gat ttt gaa gac gac gtg gag aac caa caa cag ggc ccg cag cag 2445
236 Phe Asp Phe Glu Asp Asp Val Glu Asn Gln Gln Gln Gly Pro Gln Gln
237           85           90           95
239 agg cca ccg ctg ctc cag cag tgc tgc aac gag ctc cac cag gaa gag 2493
240 Arg Pro Pro Leu Leu Gln Gln Cys Cys Asn Glu Leu His Gln Glu Glu
241       100           105           110
243 cca ctt tgc gtt tgc cca acc ttg aaa gga gca tcc aaa gcc gtt aaa 2541
244 Pro Leu Cys Val Cys Pro Thr Leu Lys Gly Ala Ser Lys Ala Val Lys
245 115           120           125           130
247 caa cag att cga caa caa cag gga caa caa atg cag gga cag cag atg 2589
248 Gln Gln Ile Arg Gln Gln Gln Gly Gln Gln Met Gln Gly Gln Gln Met
249           135           140           145
251 cag caa gtg att agc cgt atc tac cag acc gct acg cac tta cct aga 2637
252 Gln Gln Val Ile Ser Arg Ile Tyr Gln Thr Ala Thr His Leu Pro Arg
253           150           155           160
255 gct tgc aac atc agg caa gtt agc att tgc ccc ttc cag aag acc atg 2685
256 Ala Cys Asn Ile Arg Gln Val Ser Ile Cys Pro Phe Gln Lys Thr Met
257           165           170           175
259 cct ggg ccc ggc ttc tac tagattccaa acgaatatcc tcgagagtgt 2733
260 Pro Gly Pro Gly Phe Tyr
261       180
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265 tgtgttccat aaataatgta ctaatgtaat aagaactact ccgtagacgg taataaaaga 2853
267 gaagtttttt tttttactct tgctactttc ctataaagtg atgattaaca acagatacac 2913
269 caaaaagaaa acaattaatc tatattcaca atgaagcagt actagtctat tgaacatgtc 2973
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275 gtattcatta aattatgcaa gtgttctttt atttggtgaa gactctttag aagcaaagaa 3153
277 cgacaagcag taataaaaaa aacaaagttc agttttaaga tttgttattg acttattgtc 3213
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330 Asp Asp Ala Thr Asn Pro Ala Gly Pro Phe Arg Ile Pro Lys Cys Arg
331 35 40 45
333 Lys Glu Phe Gln Gln Ala Gln His Leu Lys Ala Cys Gln Gln Trp Leu
334 50 55 60
336 His Lys Gln Ala Met Gln Ser Gly Ser Gly Pro Ser Trp Thr Leu Asp
337 65 70 75 80
339 Gly Glu Phe Asp Phe Glu Asp Asp Val Glu Asn Gln Gln Gln Gly Pro
340 85 90 95
342 Gln Gln Arg Pro Pro Leu Leu Gln Gln Cys Cys Asn Glu Leu His Gln
343 100 105 110
345 Glu Glu Pro Leu Cys Val Cys Pro Thr Leu Lys Gly Ala Ser Lys Ala
346 115 120 125
348 Val Lys Gln Gln Ile Arg Gln Gln Gly Gln Gln Met Gln Gly Gln
349 130 135 140
351 Gln Met Gln Gln Val Ile Ser Arg Ile Tyr Gln Thr Ala Thr His Leu
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361 <210> SEQ ID NO: 4
362 <211> LENGTH: 2154

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/782,130

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Input Set : A:\seqlst.txt
Output Set: N:\CRF4\11042002\I782130.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; N Pos. 36³5,37¹6,37¹9,37²0,38⁰3

VERIFICATION SUMMARY

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Input Set : A:\seqlst.txt

Output Set: N:\CRF4\11042002\I782130.raw

L:293 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:3633
L:295 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:3693
L:297 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:3753